

OIPE

 RAW SEQUENCE LISTING
 DATE: 03/06/2001

 PATENT APPLICATION:
 US/09/684,016
 .
 TIME: 13:02:59

Input Set : D:\Unigene20000612.txt
Output Set: N:\CRF3\03062001\1684016.raw

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1 <1.10> APPLICANT: Kovalic, David K.
              Liu, Jingdong
      4 <120> TITLE OF INVENTION: Annotated Plant Genes 6 <130> FILE REFERENCE: 38-21(15097)D
C--> 8 <140> CURRENT APPLICATION NUMBER: US/09/684,016
     8 <141> CURRENT FILING DATE: 2000-10-10
      8 <150> PRIOR APPLICATION NUMBER: US 09/654,617
     10 <151> PRIOR FILING DATE: 2000-09-05
     12 <1.60> NUMBER OF SEQ ID NOS: 463173
     14 <210> SEQ ID NO: 1
     15 <211> LENGTH: 854
     16 <212> TYPE: DNA
     17 <213> ORGANISM: Glycine max
     19 <220> FEATURE:
     20 <221> NAME/KEY: unsure
     21 <222> LOCATION: (1)..(854)
     22 <223> OTHER INFORMATION: unsure at all n locations 24 <400> SECURENCE T
     24. <400> SEQUENCE: 1
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     28 cgtcgccgag atcggcgcgc tcctggccgc gggcaagetc gagaagatcg tcggcgtgcc
     30 cacatecaag egeacetteg ageaggegea gtegetegge atceegetet eeaegetega
                                                                             180
     32 egacaaceeg eteategace tegecatega eggtgeegac gaggtttgac eetgacetea
     34 acettgtgaa agggcggggt ggtgctcttc ttcgtgagaa gatggttgag gcagcatcgg
     36 acaagtttat tyttattytt gacgagacaa aactagttga tyggttagga gytagtgytc
     38 tagccatgcc agtggaagtt gtgcagttct gctggaagta caaccttgta agattgcagg
     40 aactgtttaa ggaggaagga gtcgaggcaa agctaaggtt tgaaggcgac aagccctatg
     42 ttactgacaa etcaaactac ategtegatt tatacttcaa gacgecaatc aaggatgegt
     44 tggcagcagg acaggaaatt gcagctctgg aaggagttgt tgaccatggg ttgttcttga
     46 acatggcgag ttcagtgatc attgctggaa cggacggtgt cagtgtcaaa acgaagtgag
                                                                             660
     48 tittigagit gettigtigt tigtgitigag tittlacaatt taatgiacaa getathggig
                                                                             720
     50 taaaagcage tgataagatg etteaataaa ggtgtteece eaatgtgggg attgaetate
W--> 54 nnnnnnnnn nnnn
     57 <210> SEQ ID NO: 2
     58 <211> LENGTH: 1,222
     59 <212> TYPE: DNA
     60 <213> ORGANISM: Glycine max
     62 <400> SEQUENCE: 2
     64 acgcgtagto totatotott ttotocagoa goaccaaaac cacccaacac togaactcaa
     66 tatccattat taccaaaaaa aaagaaaaaa ggatatcaaa caccetttte atcattaaaa
                                                                             1.20
     68 eteteagite ectetitece tateaaaace etiteeeaca acaetactea etecagitaca
    70 agaaaccett titeaatiig gitteaacag cacacacac cacacatata tatatatata 72 gealgittae eitgaaleat tetteigati igiaccatgi ticecetgag eleteateti
     74 cettggacae atectogoog getteggagg getetegtgg egtggeattt teegaegagg
     76 aggtgegget ggeggtgagg caccegaaga agegggeagg teggaagaag tteegggaga
     78 cgcgccaccc qgtgtaccqq qqqqtqaqqa qqaqqaactc qqataaqtqq gtqtqtqaqq
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80 tgagggagcc caacaagaag accaggattt ggctggggac tttccccacg ccggagatgg

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82 cggctcgggc gcacgacgtg gcggcaatgg ccctgagggg ccggtatgcc tgtctaaact 600
    84 ttgctgactc ggcctggcgg ttacctgttc ccgccacggc cgaggcaaag gatatacaga 660
    86 aggeageage agaagetgee eaggetttea gaceagatea aacettaaaa aatgetaata
    88 caaqqoagga qtqtqtqqaq qcqqtqqcqq tggcggtggc ggagacaaca acqqcqacqq
     90 cacaagyggt gttttatatg gaggaagaag agcaggtgtt ggatatgcct gagttgetta
    92 ggaatatggt gctcatgtcc ccaacacatt gcttagggta tgagtatgaa gatgctgact
    94 tggatgecea ayatgetgaq gtgteaetat ggaatttete aatttaataa tgtgttttgg
     96 tittggttitt gitgttagtt tittggagty aacagigtei glaciggitt titattagia 1020
    98 gtacggatac tagttatagg tggaaagatt gcgagggacc aaaaggaatt ttcttttgaa 1080
    100 cccttttttg tcaatgtaat caatatcacg tatatcatga agtgaatccc ttcaagttta 1140
    102 tgtataaatt aaataaaaga aaaatctagt tttggttatg aatctttgtt aaagtaaaaa 1200
    104 aaaaaaaaa aagggeggee ge
    107 <210> SEQ ID NO: 3
    108 <211> LENGTH: 708
    109 <212> TYPE: DNA
    110 <213> ORGANISM: Glycine max
    112 <400> SEQUENCE: 3
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    116 ggcacgggcc cacgacgtgg cggcaatggc cctgagggyc cggtatgcct gtctcaactt
    118 egetgacteg aegtggeggt taccaattee egecaetget aaegcaaagg atatacagaa 180
    120 agcagcagca gaggetgeeg aggettteag accaagteag acettagaaa atacgaatae
    122 aaagcaagag tgtgtaaaag tggtgacgac aacaacgatc acagaacaaa aacgaggaat
    124 gttttatacg gaggaagaag agcaagtgtt agatatgcct gagttgctta ggaatatggt
    126 gottatytee ceaacacatt geatagggta tgagtatgaa gatgetgaet tggatgetea
                                                                        420
    128 agatgetgag gtgteectat ggagtttete aatttaataa egtgettttg gtttggtttt 480
    130 ttatgttagt ttttggagtgt gactgtctgt actggttttt tattagtagt acggatacta
    132 getataggtg geagattgaa agggaceaaa aggaatttte ttttgaaace ettttgtca
    134 aagtaatcaa tegegtatea teaagtgaat eeettgatea agtttatgta tgaattaaat 660
    136 aaaagaagaa totagttttg gtaaaaaaaa aaaaaaaagg geggeege
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    139 <210> SEQ ID NO: 4
    140 <211> LENGTH: 551
    141 <212> TYPE: DNA
    142 <213> ORGANISM: Glycine max
    144 <220> FEATURE:
    145 <221> NAME/KEY: unsure
    146 <222> LOCATION: (1)..(551)
    147 <223> OTHER INFORMATION: unsure at all n locations
    149 <400> SEQUENCE: 4
    151 contrattoa ctototttto teeggeagea ceaaaaccae ccaacacttg aaateaatat
    153 ccattgttac caaaaaagaa agatcaaaac accetttaca tcattatact cagtteectc 120
    157 ctaactttgg cttcagcggc acatatatat atatatata atagaacatg tataccttga
    159 accaetette atattigtae eatgitteee etgagetete etetteetig gaetegieet 300
    161 egeoggette egagggetet egeggegtgg catttteega egaggaggtg eggetggegg
    163 tgaggcaccc gaagaagcgg geegggegga agaagtteeg ggagaegege caceeggtgt
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    165 tecegggegt gtggaggatg gtettggtte tittgggttte titatitata attatigati 480
W--> 169 nnnnnnnnn n
    172 <210> SEQ ID NO: 5
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Input Set : D:\Unigene20000612.txt
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	<211> LENGTH: 972 . <212> TYPE: DNA											
	<213> ORGANISM: Glyci	ne may										
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	acattttcaa atccccactg					120						
	cagaageate ttattegtet					180						
	cgtcagcgcg gccaaagaag					240						
	totacogagg agtgcgtcgc					300						
	acaacaactc acggatttgg					360						
	acgacgttgc ggcgctcgcg					420						
	ggtggcggct gacggtgccg					480						
	aggetgetga ggeatttgea					540						
	gtgtgatggc cacgaatgat			-		600						
	acttgcatga tttgcttttg					660						
	caagagatgg tagggactgg					720						
	ggaacttete aatttgaege					780						
	aagegaaaae acctttttgt					840						
	gataaattaa gtagcactcg					900						
	totatgattt gtttcgtgat					960						
	aaqqqqqqc qc	- Caua Coocaa	.,	4900990444		972						
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	<211> LENGTH: 862											
	<212> TYPE: DNA											
	<213> ORGANISM: Glycin	ne max										
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	gacgacggcg gccatggcga					120						
	caagacggtg taccttgtgg					180						
	cttccgctgc caccactgca			-		240						
	agtgetetae tgeaggeete					300						
	gagettegaa ggaacteega					360						
233	tactaaagtc tcaagcgcct	ttgctggcac	cagagagaaa	tgtgttggat	gcagcaagac	420						
235	agtetateca actgagaggg	tcactgttaa	taacactatg	taccacaaga	gctgcttcaa	480						
237	gtgct.gccat ggagggtgca	ccatcagece	ttctaactac	attgcgcacg	aggggaaget	540						
239	gtactgcaag caccaccaca	tccagctgat	caaggagaag	gggaacttca	gccagcttga	600						
241	gaatgaccac gagaagacat	cacaggetgg	gtcactggag	gaggatgaac	aagagtattg	660						
243	atcactgage acaaccacag	atgaataatg	aatcctcttg	catttgccta	gagcactatg	720						
245	tatttctgtg gcgtggttcg	atttcagttt	taccaatgga	ggtcatgtgt	gtt.cagagaa	780						
247	caatgatatt gttgtcttgt	atctgcatgt	aaacgttgtt	atctttgagt	cagaaaaaaa	840						
249	aaaaaaaaa aagggcggcc	gc				862						
252	<21.0> SEQ ID NO: 7			•								
253	<211> LENGTH: 1686											
254	<21.2> TYPE: DNA											
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	theetttigg thetetiggt					120						
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265 gctaatgget ttaatgeeta ttggetaatg tatettgeet etgateette teagagaaac 240
267 adagteteat cagtitttea acaggettea aateatggee teaacattge cagaacttgg
269 gettteagtg atggtggata teaaceacta caatactete etggatetta caattaceaa
271 atgtttcagg gcttggattt cgcaatagcc gaagcaagaa aatatgggat caagatggtg 420
273 ttgagtttgg tgaataacta tgagaacatg ggtggaaaga aacagtatgt ggaatgggca
275 aggagteagg gacagteeat aaattetgag gatgaetttt ttacaaaatt eggetegagg
                                                                       540
277 ggatactaca aaaaccacat caaggotgta ottacaagac gtaatagcat cactggagtt
279 gettacaaag acgacccaac tataatgget tqqqaactta tqaatqagat taqqtqccct
281 totgateaat caggaaggac agtteagget tggateactg agatggeate ttacctgaaa
283 tocatagatg gaaaccactt gotggaagot ggtotggagg gtttotacgg ocagtoaaag
285 cotcagtota acceaaactt caatgtagga accgatttea ttgccaacaa ccaaateeca
287 ggcattgatt tegeaacagt gcaetectae eeggaceaat ggctateaag ttetagttat
289 gaggaccaaa teteatttt aggeegttyg etggatgage acateeaaga tyeacagaac
291 accetteaca agecaettet attigeggag titiggtatti ccacaaagag tiatggtggc 1020
293 aactcaacac caagggateg gttattcaac acagtatatt etgcaatata etcatcagca 1080
295 ageagtggeg gggetgetgt tggtggettg ttttggcaac ttatggttca aggaatggat 1140
297 tettategag aeggttaega agttgtetta gatgagagee etteaaegge taatttgate 1200
299 gctcaagagt ctcaaaaaact aaaccgaatt cgcaagatgt acgccagact tagaaacatt 1260
301 gagaagtgga atgaagctaa gcaaattaga ggtggaaact gaatatgata cgtcttaact 1320
303 accatgatat atatatatag agagagagag agagagtgaa gggagttaga aactttaatt 1380
305 gataacttat tigggtgtgg attaatette agaatgaagt gicccaataa tettagtgge 1440
307 aagaattgoo tgttattitto gtoaaagtat tooaagtgaa gtgttotaca aatqogatag 1500
309 acataactat ttattattga gatatttega gagaaacttt gtaatgeaeg aactageatt 1560
311 teatetacae tattatgteg geaagtgaat gtactttgte taaactgttt ataaaatgga 1620
313 ctyctgacga taatttgaag tycttcaaaa aaaaaaaaaa aaaaaaaaa aaaagggcyg 1680
315 ccgccg
318 <210> SEQ ID NO: 8
319 <211> LENGTH: 1661
320 <212> TYPE: DNA
321 <213> ORGANISM: Glycine max
323 <400> SEQUENCE: 8
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327 ggagtttygt atttccacaa agagttatgg tggcaactca acaccaaggg atcggttatt 120
329 caacacggta tattcagcaa tatactcatc agcaagcagt ggcggggctg ctgttggtgg
331 cotgetting caactuateg cicaaggaat ggattottat ogagacgett acgaggingt
333 cttagatyag agccettcaa eggetaattt gattgeteaa gagteteaaa aactaaaceg
335 aattogoaag atgtacgooa gactoagaaa cattgagaaa tggaatgaag otaagoaaat
337 tagaggatga aactgaatat atatatata atatagagag agagagagag agagaagggt
339 gagttagaaa etttaattta taaettaate gtagtggcaa gaattgeetg ttattacaag
341 tgaagtgtoo tacaaatgca ataaacataa ctattataga gatatatega gagaaacttt
                                                                       540
343 gtaatgtacg agctagcatt tcatctacta ttatgtcgga agtgaatgta cttatgtcaa
345 aactattata aaytggcctg ctggataact tgaaqtgctt catataactg ttccatcttt
347 gattgttgca ttttttaåtg taactgtgat geoogaecta atgtagaate etteetttge
349 cetyteaget teltgaacae categytytt tyctatgate acatttette etatettyge
351 attettgteg attatgeaat teetgatttt agtatttgee eegacaceaa ttggaacett
                                                                      840
353 coeffective accagagaty caattteata etcagtftga taatagteag cacceateat
355 categratee tgaageteea caecagaete caaaegtgag egtaeteeaa caatagaatg
357 tigaacqciq cacteccica agaaqcaacc atqaqatata attqcateca caatciiqeq 1,020
359 gttttctact ttggtaggtg gtaggaatct gggggaagtg aagaaaggtg tctttggatc 1080
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Input Set : D:\Unigene20000612.txt
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361 atagaattea aatttaggag getgttetgt tagageaaga tttgeateaa agaaggaett 1140
363 tatagtteea atatetteee agtagteatt gaacaaatat geetggacat tgtgeteatt 1200
365 cacageagat gggataattt cagateeaaa qteattgeat gaagaacatt tecatettaa 1260
367 tagttqcaqc aaqqtttcaq ttctaaacac qtaqacaccc atqqatqcaa tataaqqatq 1320
369 tittitetget tettgitggea ataaccetaa aagagtggitg teaacacgea tigeettiag 1380
371 atotgatoco ttaggttttt otgoaaactg tataatoogt cotgtottat caattttoat 1440
373 cagtocatag totgatgood gactgteato catgggtaca catgaaactý tgatatoago 1500
375 atttgtgtca acatgtctct gtacaaagtt catgtagtcc attcggtaaa gatgatcacc 1560
377 agaaagtate aatatatget caacattett gttettggea teetcaaaaa eecatataaa 1620
379 ttgtcttaca gcatcggcgg tcccttggaa ccacttcttc t
382 <210> SEQ ID NO: 9
383 <211> LENGTH: 407
384 <212> TYPE: DNA
385 <213> ORGANISM: Glycine max
387 <400> SEQUENCE: 9
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391 gtttttgaga etettgageg ateaaattag eegttgaagg geteteatet aagaeaaett
393 ogtaacogto togataagat gtacgccaga ottagaaaca ttgagaagtg gaatgaagot
                                                                        180
395 aagcaaatta gaggtggaaa ctgaatatga tacgtettaa etaccatgat atatatatat
                                                                        240
397 agagagagag agagagagt aagggagtta gaaactttaa ttgataactt atttgcgtgt
399 ggattaatet teagaatgaa gtgteecaat aatettagtg geaagaattg eetgttattt
                                                                        360
401 tegteaaaaa aaaaaaaaaa aaaaaaaaaa aagggeggee aeegeet
404 <210> SEQ ID NO: 10
405 <211> LENGTH: 614
406 <212> TYPE: DNA
407 <213> ORGANISM: Glycine max
409 <400> SEQUENCE: 10
411 ctttaccatt etccatggag tggeggggaa ttattatgtt tatactataa tataataaca
413 taattatatt tatagtaact actacttatt tattttttta aaatatctta atttcccacc
415 ttaccetaca atcaaggaca gaatetatge aatttaaaag ecaacaacgt tgagagaage
417 caagtgtgaa gacagacagt gagaaatgat gatgaagegt eggtteettt tggttetett
                                                                       240
419 ggttctctat gtcacagtgg agcagagtaa gctactccac cacgtagaag ccgatggtgg
                                                                        300
421 gtttgtaaaa acaagaggag tgcagcttat gctgaatggg agtccctact atgctaatgg
423 etttaatgee tattggetta tgtaeettge etetgateet teecagagaa acaaaatete
425 atoggtattt caacaggett caaatcatgg actcaacatt geeagaactt gggeetteag
                                                                        480
427 tgatggtgga tatcaacccc tacaatactc tcctggatcc tacaatgacc aaatgtttca
                                                                        540
429 gggcttggat ttcgcaatag ccgaagcaag aaaatatggg atcaagatgg tgttcgagtt
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431 ggtgaataac tatg
434 <210> SEQ ID NO: 11
435 <211> LENGTH: 1587
436 <212> TYPE: DNA
437 <213> ORGANISM: Glycine max
439 <400> SEQUENCE: 11
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443 taagtttett gtttgattag aatggageee caacacteet ecatetacaa tettacaaac
445 ctitattega agttetgatg getgtgegat cattetetta egatgtgtte etcagettee
447 gaggggaaga tactogttat ggtttoactg gctatotota caatgtoott ogggaaaggg
449 gaattcacac etteattgat gaegaegage eecaggaagg ggaegaaate aegaeageae
                                                                       300
451 ttgaggagge tattgagaag tecaagattt teateategt getetetgaa aactaegeat
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## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

 VERIFICATION SUMMARY
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Input Set : D:\Unigene20000612.txt
Output Set: N:\CRF3\03062001\1684016.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application No L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  $\rm L\!:\!167~M\!:\!341~W\!:$  (46) "n" or "Xaa" used, for SEQ ID#:4 L:169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 L:562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 L:693 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 L:722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 L:739 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 L:1266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31L:1268 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 L:1270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 L:1453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 L:1778 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 L:1780 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45L:1782 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 L:1873 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 L:1926 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 L:1928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 L:2947 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 L:2949 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 L:3286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 L:3329 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 L:3331 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 L:4094 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 L:4151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:96 L:4994 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 L:4996 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 L:5459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:123 L:5461 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:123 L:5654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127 L:5755 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:130 L:5757 M:341 W: (46) "n" or L:6154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:130 "Xaa" used, for SEQ ID#:139 L:6397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:144 L:6403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:144 L:7218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:162 L:7361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:165 "Xaa" used, for SEQ ID#:173 L:7776 M:341 W: (46) "n" or L:7877 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175 L:7879 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175 L:7881 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175 L:7928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176 L:7930 M:341 W: (46) "n" or "Xaa" used, for SEQ TD#:176 L:7932 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176 L:7977 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177

L:8508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:185

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/684,016

DATE: 03/06/2001 TIME: 13:03:00

Input Set : D:\Unigene20000612.txt
Output Set: N:\CRF3\03062001\1684016.raw

L:8510	M:341	W:	(46)	" n "	or	"Xaa"	used,	for	SEQ	ID#:185
L:8603	M:341	W :	(46)	u U u	or	"Xaa"	used,	for	SEQ	TD#:187
L:8605	M:341	W :	(46)	" II "	o.r	"Xaa"	used,	for	SEQ	TD#:187
L:8768	M:341	W :	(46)	$^{n}$ II $^{n}$	or	"Xaa"	used,	for	SEQ	ID#:192